

FIG. 2

Consensus	MALPHILF TVLSPFILT APPPCOMTS SSPYQEFAR WOPGCHIDAP SYRSLSKGTP TFIATHTMR NCHNSATLCH HANTHTWTK MINPSCFGGL	100
P Env DNA 10	.....	100
P Env DNA 21	.....	100
P Env DNA 6	.....	100
Consensus	GTVVCTTFT QTKMSGGGV QDQREBVK EVISQURVH GTSSTPGLD LSKLHETLT HTRVLSLNT TLTGLHNSA QNFTNCWICL PLNFRPYVSI	200
P Env DNA 10	.....	200
P Env DNA 21	.....A.....	200
P Env DNA 6	.....	200
Consensus	FVPEQNNFS TEINTTSVLV GLVSNLEIT HSNLTCVK SNTYTINSQ CINWTPPTQ IVCLPSGIFF VQTSATRCI NGSSSHCFL SELVPPHTIY	300
P Env DNA 10	.....	300
P Env DNA 21	.....A.....	300
P Env DNA 6	.....	300
Consensus	TEQDLYSVI SKPRNRUPI LPFVIGAGVL GALTGIGCI TTSTQFYKL SQELNGWER VADSLVTLOD QIASLAAVL QNRALDILT AERGCTCLFL	400
P Env DNA 10	.....N.....	400
P Env DNA 21	.....	400
P Env DNA 6	.....	400
Consensus	GECCYTVNQ SOIVTERVKE IROIRIGRAE ELRWTFMGL LSQNHMILP FLOPLAAILL ULLEFGCIEN LWNFVSSRI EAVKLQMEK HOSKTKLYRR	500
P Env DNA 10	.....	500
P Env DNA 21	.....	500
P Env DNA 6	.....	500
Consensus	FLDRPASPRS DVNDINGTTP EEISAAQPLL RENSAGSS	538
P Env DNA 10	.....	538
P Env DNA 21	.....	538
P Env DNA 6	.....	538

FIG. 2

Consensus	120
Env DNA 6	120
Env DNA 10	120
Env DNA 21	120
Consensus	240
Env DNA 6	240
Env DNA 10	240
Env DNA 21	240
Consensus	360
Env DNA 6	360
Env DNA 10	360
Env DNA 21	360
Consensus	480
Env DNA 6	480
Env DNA 10	480
Env DNA 21	480
Consensus	600
Env DNA 6	600
Env DNA 10	600
Env DNA 21	600
Consensus	720
Env DNA 6	720
Env DNA 10	720
Env DNA 21	720
Consensus	840
Env DNA 6	840
Env DNA 10	840
Env DNA 21	840
Consensus	960
Env DNA 6	960
Env DNA 10	960
Env DNA 21	960
Consensus	1080
Env DNA 6	1080
Env DNA 10	1080
Env DNA 21	1080
Consensus	1200
Env DNA 6	1200
Env DNA 10	1200
Env DNA 21	1200
Consensus	1320
Env DNA 6	1320
Env DNA 10	1320
Env DNA 21	1320

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FIG. 2 continued

Consensus	CTGAGCGAAT GAGTGGCTG GATTCGCGC TCTGAGAC CTGAGGAG TATATATG CAGCTGCTT TGGAGCTG TACTTTAC CTGCTTGTA ACTTGTCTC TTCCAGATC	1440
Env DNA 6	.....	1440
Env DNA 10	.....	1440
Env DNA 21	.....	1440
Consensus	CAAGCTGTA AACGCAAT GAGGCGAG ATGGAGTGA AGCTAGAT CTAGCGGA CCGCTGAG CCGCTGCTG CCGAGAT CATGTATG ACATCAAGG CAGCCTCTCT	1560
Env DNA 6	.....	1560
Env DNA 10	.....	1560
Env DNA 21	.....	1560
Consensus	GAGGAAATC CAGCTGCA AGCTGACTA CCGCGAAT CAGCGAG CATTAG	1617
Env DNA 6	.....	1617
Env DNA 10	.....	1617
Env DNA 21	.....	1617

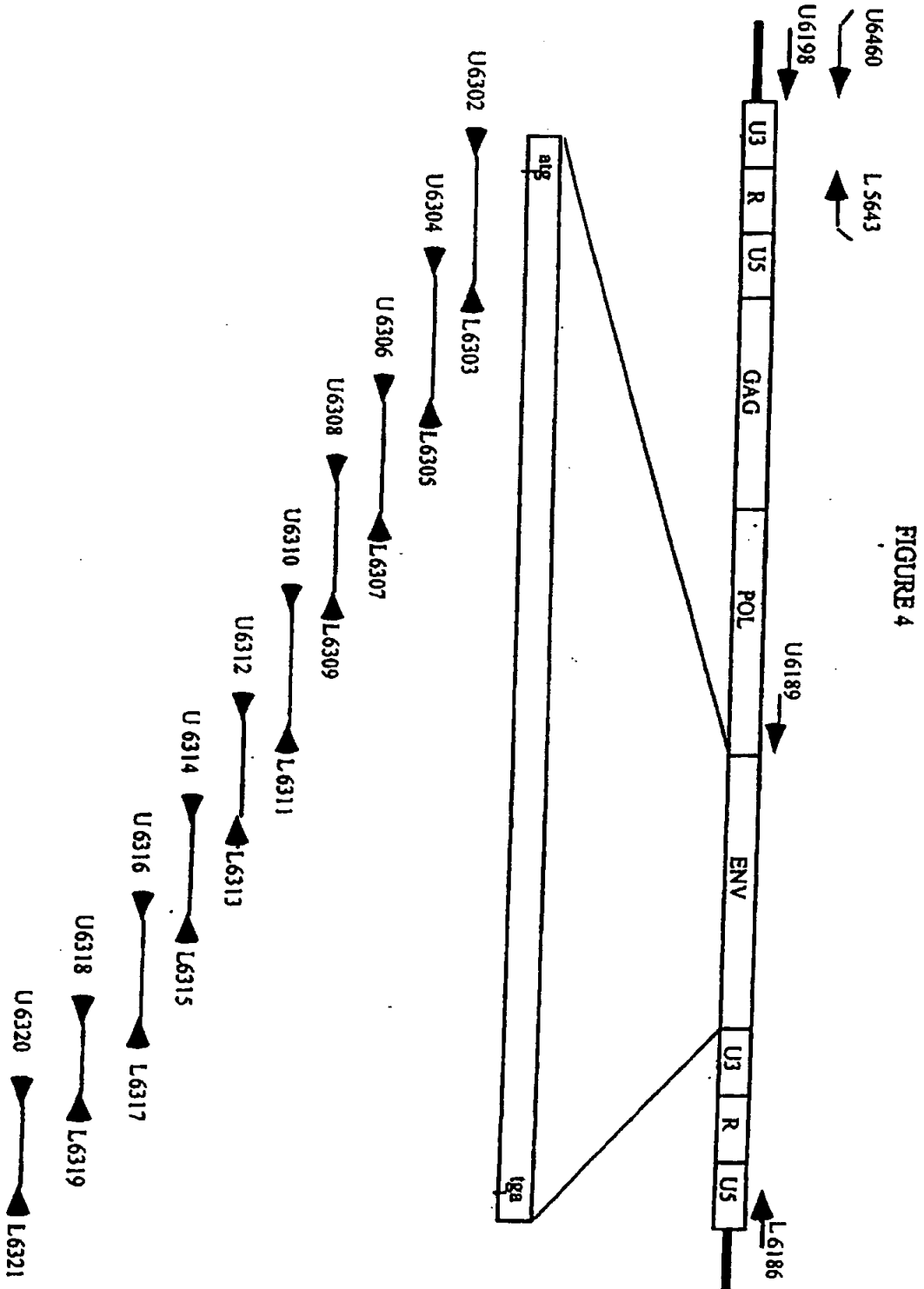
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FIG. 3

Consensus	TCAGGACAG GACTAGCTGG ATTGCTTAGG CCGACTTAGA ATCCCTTAGC CTAGCTGGG ARGTCACAC GTCCACTTT AACACGGGG CTTCGACTT	100
LTR6 c1A	.....	100
LTR21 c1S	.....G.....A.....	100
Consensus	AGTCACACC TCACCAATCA GAGGCTTCAC TAAATGCTA ATTAGGAAA GACGGAGGT AAGAAATAG CCAATCACT ATTGCTTAGG ACCACAGGAG	200
LTR6 c1A	.....	200
LTR21 c1S	.....A.....G.....	200
Consensus	GAGGACAAV PATCGGATA TAAACCGAG YMTTCGACTY GGCAC	246
LTR6 c1A	.....T G.....A. TC.....C.....	246
LTR21 c1S	.....C A.....G. CA.....T.....	246

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FIG. 4



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FIG. 5

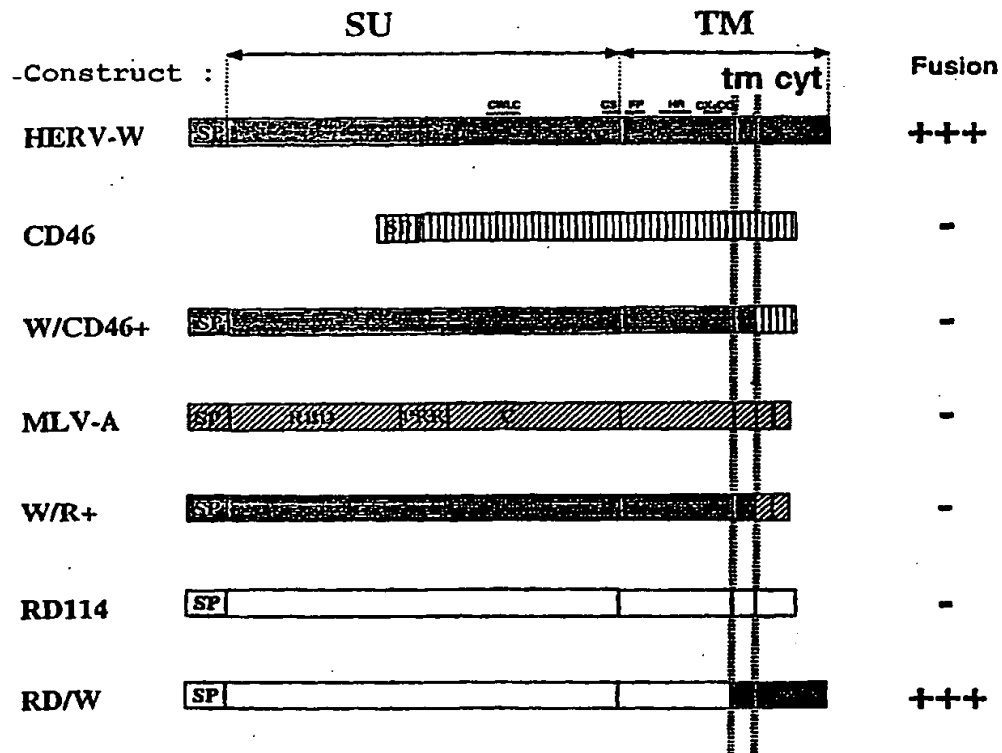


Figure XXX [sic]. Scheme and characterization of the chimeric Env HERV-Ws